## SEQUENCE LISTING

(1) GENERAL INFORMATION:

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- (i) APPLICANT: Olsen, et al.
- (ii) TITLE OF INVENTION: Vascular Endothelial Growth

Factor 3

- (iii) NUMBER OF SEQUENCES:
  - (iv) CORRESPONDENCE ADDRESS:
    - (A) ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN,

CECCHI, STEWART & OLSTEIN

- (B) STREET:\ 6 BECKER FARM ROAD
- (C) CITY: \ROSELAND
- (D) STATE: NEW JERSEY
- (E) COUNTRY: USA
- (F) ZIP: 0∀068
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: \ 3.5 INCH DISKETTE
  - (B) COMPUTER: IBM PS/2
  - (C) OPERATING SYSTEM: MS-DOS
  - (D) SOFTWARE: WORD PERFECT 5.1
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER:
  - (B) FILING DATE: condurrently
  - (C) CLASSIFICATION:
- (vii) ATTORNEY/AGENT INFORMATION:
  - (A) NAME: FERRARO, GREGORY D.
  - (B) REGISTRATION NUMBER: \36,134
  - (C) REFERENCE/DOCKET NUMBER: 325800-
- (viii) TELECOMMUNICATION INFORMATION:
  - (A) TELEPHONE: 201-994-1700
  - (B) TELEFAX: 201-994-1744

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- (2) INFORMATION FOR SEQ ID NO:1:
- (i) SEQUENCE CHARACTERISTICS
  - (A) LENGTH: 666 BASE PAIRS
  - (B) TYPE! NUCLEIC ACID
  - (C) STRANDEDNESS: SINGLE
  - (D) TOPOLOGY: LINEAR
- (ii) MOLECULE TYPE: cDNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

60 ATGAGAAGGT GTAGAATAAG TGGGAGGCCC CCGGCGCCCC CCGGTGTCCC CGCCCAGGCC CCTGTCTCCC AGCCTGATGC CCTGGCCAC CAGAGGAAAG TGGTGTCATG GATAGATGTG 120 TATACTCGCG CTACCTGCCA GCCCCGGGAG GTGGTGGTGC CCTTGACTGT GGAGCTCATG 180 GGCACCGTGG CCAAACAGCT GGTGCCCAGC TGCGTGACTG TGCAGCGCTG TGGTGGCTGC 240 TGCCCTGACG ATGGCCTGGA GTGTGTGCCC ACTGGGCAGC ACCAAGTCCG GATGCAGATC 300 CTCATGATCC GGTACCCGAG CAGTCAGCTG GGGGAGATGT CCCTGGAAGA ACACAGCCAG 360 TGTGAATGCA GACCTAAAAA AAAGGACAGT GCTGTGAAGC CAGACAGGGC TGCTACTCCC 420 CACCACCGTC CCCAGCCCCG TTCTGTTCCG GGCTGGGACT CTGCCCCCGG AGCACCCTCC 480 CCAGCTGACA TCACCCAATC CCACTCCAGC CCCAGGCCCC TCTGCCCACG CTGCACCCAG 540 600 CACCACCAGT GCCCTGACCC CCGGACCTGC CGCTGCCGCT GTCGACGCCG CAGCTTCCTC CGTTGTCAAG GGCGGGGCTT AGAGGTCAAC CCAGACACCT GCAGGTGCCG GAAGCTGCGA 660 666 AGGTGA

- (2) INFORMATION FOR \$EQ ID NO:2:
  - (i) SEQUENCE CHARACTERISTICS
    - (A) LENGTH: 221 AMINO ACIDS
    - (B) TYPE: AMINO ACID
    - (C) STRANDEDNESS:
    - (D) TOPOLOGY: LINEAR
  - (ii) MOLECULE TYPE: | PROTEIN

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## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Met Arg Arg Cys Arg Ile Ser Gly Arg Pro Pro Ala Pro Pro Gly Val Pro Ala Glh Ala Pro Val Ser Gln Pro Asp Ala Pro Gly His Gln Arg Lys Val\Val Ser Trp Ile Asp Val Tyr Thr Arg Ala Thr Cys Gln Pro Arg Glu Val Val Val Pro Leu Thr Val Glu Leu Met Gly Thr Val Ala L $\chi$ s Gln Leu Val Pro Ser Cys Val Thr Val Gln Arg Cys Gly Gly Cys Cys Pro Asp Asp Gly Leu Glu Cys Val Pro Thr Gly Gln His Gln Val Arg Met Gln Ile Leu Met Ile Arg Tyr Pro Ser Ser Gln Leu Gly Glu Met Ser Leu Glu Glu His Ser Gln Cys Glu Cys Arg Pro Lys Lys Lys Asp Ser Ala Val Lys Pro Asp Arg Ala Ala Thr Pro His His Arg Pro Gln Pro Arg Ser Val Pro Gly Trp Asp Ser Ala Pro Gly Ala Pro Ser Pro Ala Asp Ile Thr Gln Ser His Ser Ser Pro Arg Pro Leu Cys Pro Arg Cys Thr Gln His His Gln Cys Pro Asp Pro Arg Thr Cys Arg Cys Arg Cys Arg Arg Arg Ser Phe Leu Arg Cys\Gln Gly Arg Gly Leu Glu Leu Asn Pro Asp Thr Cys Arg Cys Arg Lys Leu Arg Arg 

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